

Cockrell Creek *TMDL for Shellfish Waters*

Heathsville, VA



5 September 2007

What is a TMDL?

TMDL = Total Maximum Daily Load =
maximum amount of a pollutant that can
enter a waterbody without violating water
quality standards (WQS)

WQS = numeric or narrative limits on
pollutants that ensure the protection of
human health and of aquatic life

The TMDL Process in VA

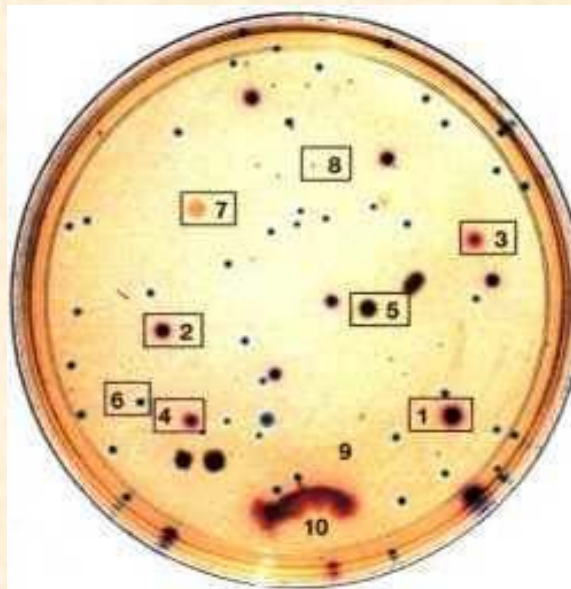
- Three Step TMDL Process in VA
 - TMDL Development (Find the source of the pollutant & determine the reduction needed)
 - Implementation Plan Development (Identify conservation measures to fix the problem. Conservation measures are often called Best Management Practices or BMPs.)
 - Implement the BMPs and sample to see improvement.

Why a TMDL for the Cockrell Creek Watershed?

- VDH Division of Shellfish Sanitation (DSS) monitors fecal coliform levels in shellfish waters
- Applicable water quality standard: 30-month geometric mean not exceeding 14 MPN/100 mL, and a 90th percentile not exceeding 49 MPN/100 mL
- Cockrell Creek has observed exceedances that necessitate TMDLs be developed to bring them into compliance

What are fecal coliform bacteria?

- Bacteria present in the intestines of warm blooded animals, like livestock, wildlife, birds, and humans
- Indicator of the potential presence of pathogens in water



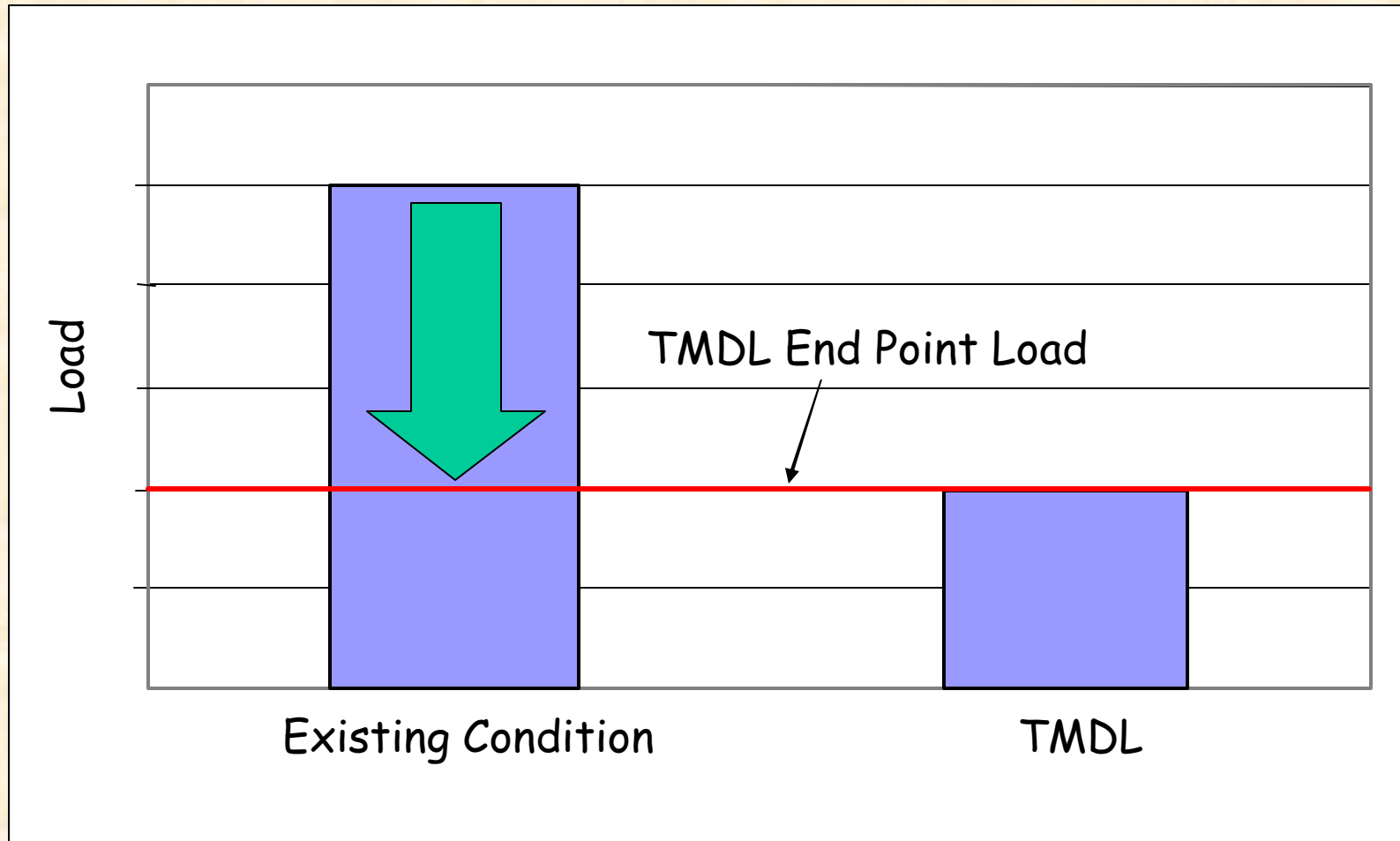
Required Elements of a TMDL

- Developed for critical stream conditions;
- Must meet water quality standards;
- Consider seasonal variations;
- Include wasteload and load allocations;
 - waste load = permitted human discharges
 - load allocation = loads due to wildlife, cattle, pets
- Include a margin of safety;
- Consider impacts of background contributions;
- Be subjected to public participation;
- Have reasonable assurance for implementation.

What information might be used to develop a TMDL?

- VDH Sanitary Shoreline Survey
- VDH Bacteria monitoring
- Population estimates (human,pets,livestock)
- Affected waters volume
- Bacterial Source Tracking Data
- Land use, Climate, Tide and similar data
- DEQ permit data if applicable
- DEQ spill response and remediation data

An Example TMDL



Reducing existing bacteria load to the TMDL end point load is expected to restore water quality.

What's changed since last year?

- New data
 - DEQ Special Study conducted from Aug 2006 – Feb. 2007
 - Revised data analysis
 - New studies and information incorporated into TMDL.

COCKRELL CREEK

➕ Reedville STP Outfall

■ Omega Protein

DH Condemnation Zones

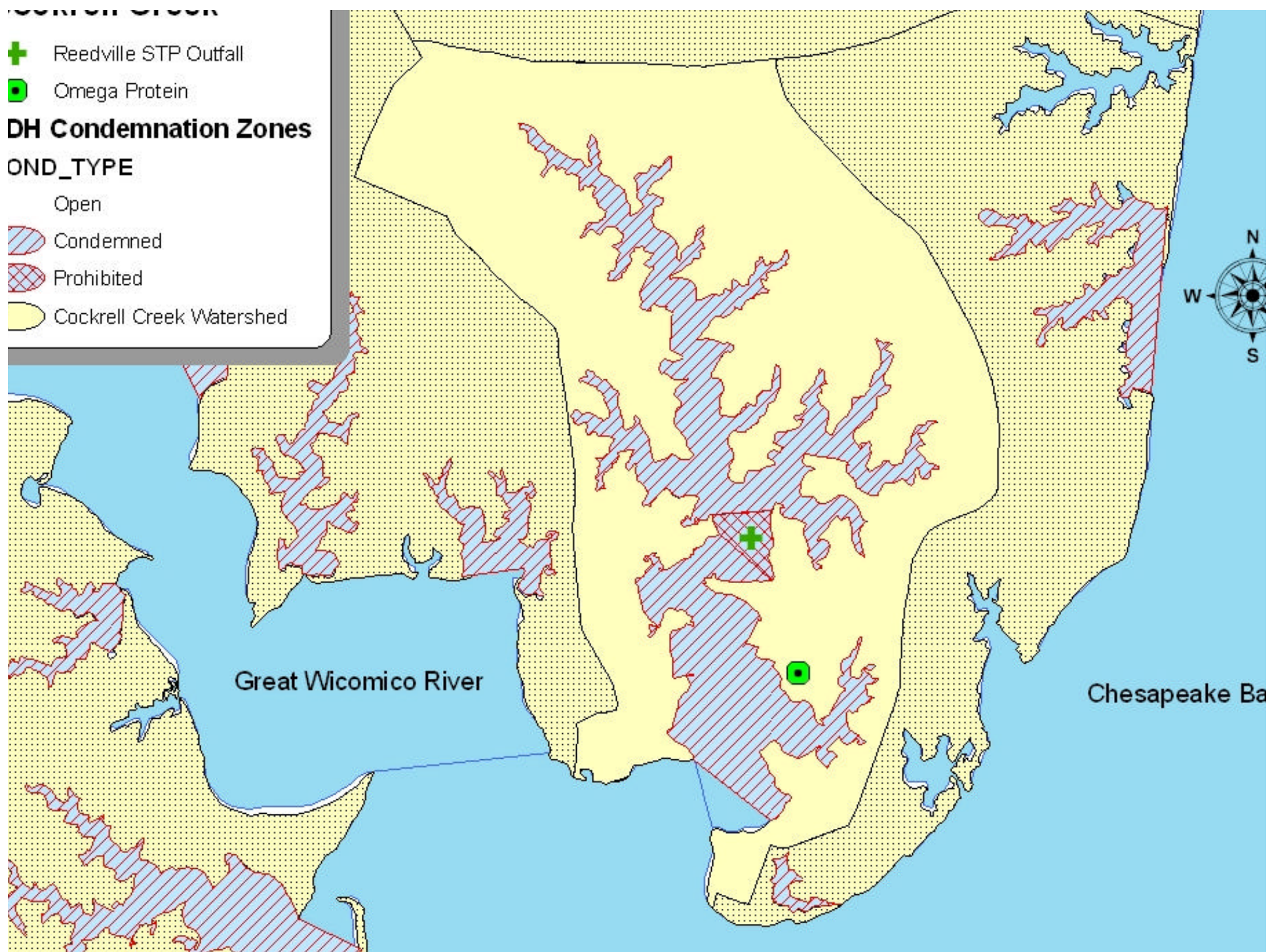
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Open

Condemned

Prohibited

Cockrell Creek Watershed



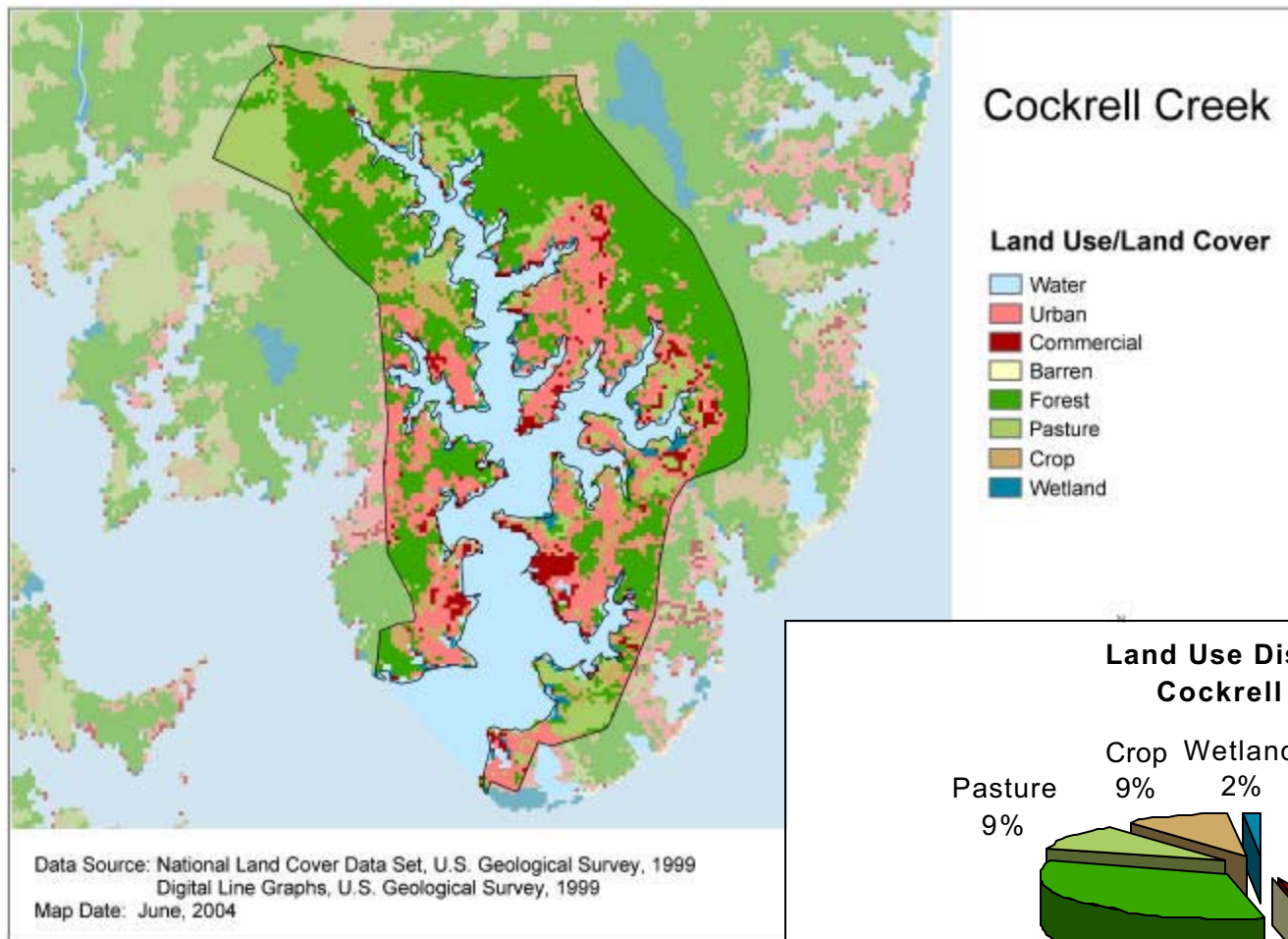
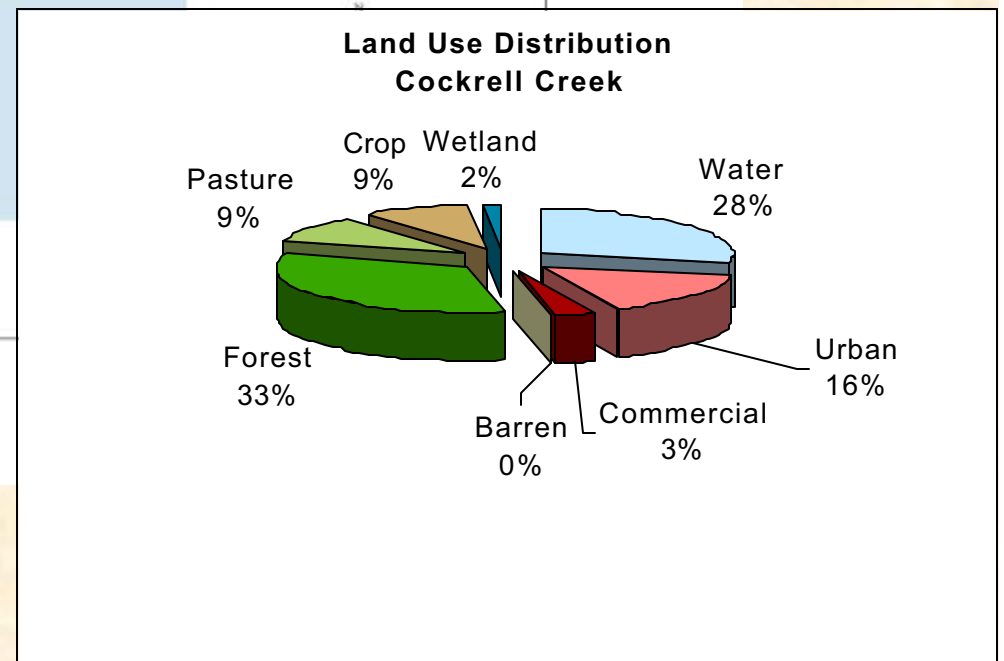


Figure 3.1



**Table 3-1 Animal Populations and Septic Systems
Growing Area 12**

Fecal Coliform Sources	2 Cockrell Creek
Cattle	21*
Horse	6*
Pig	0
Deer	38
Duck	485
Geese	334
Raccoon	85
Dog	55
Estimated Septic Systems	90

* - from DSS Shoreline survey 2004.

Estimated Human population = 392. (2000 U.S. Census)

Approx. 60% of urban area with public sewer. (Analyzed via GIS data from NNPDC.)

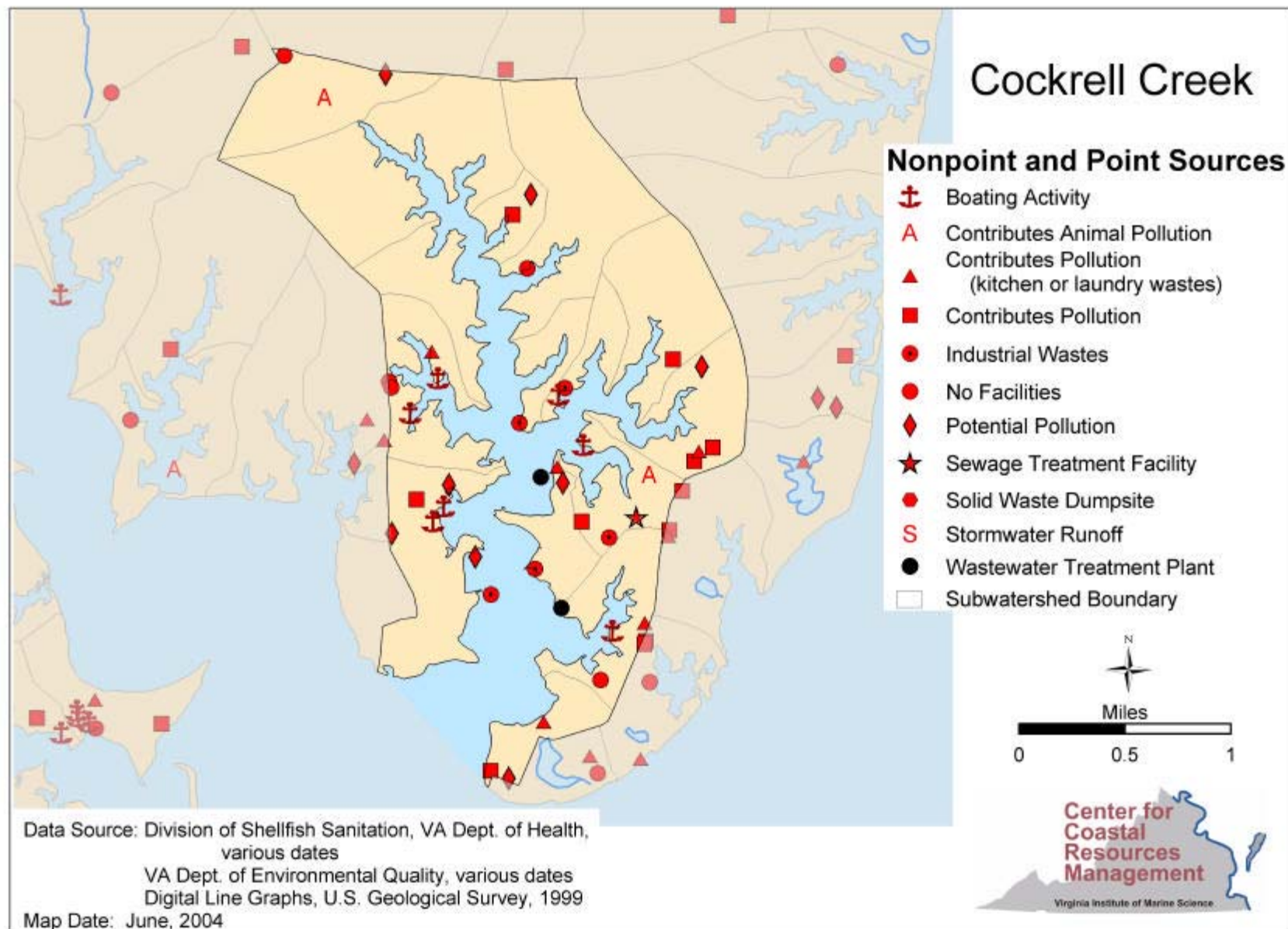


Figure 4.4

Water Quality Monitoring & Data

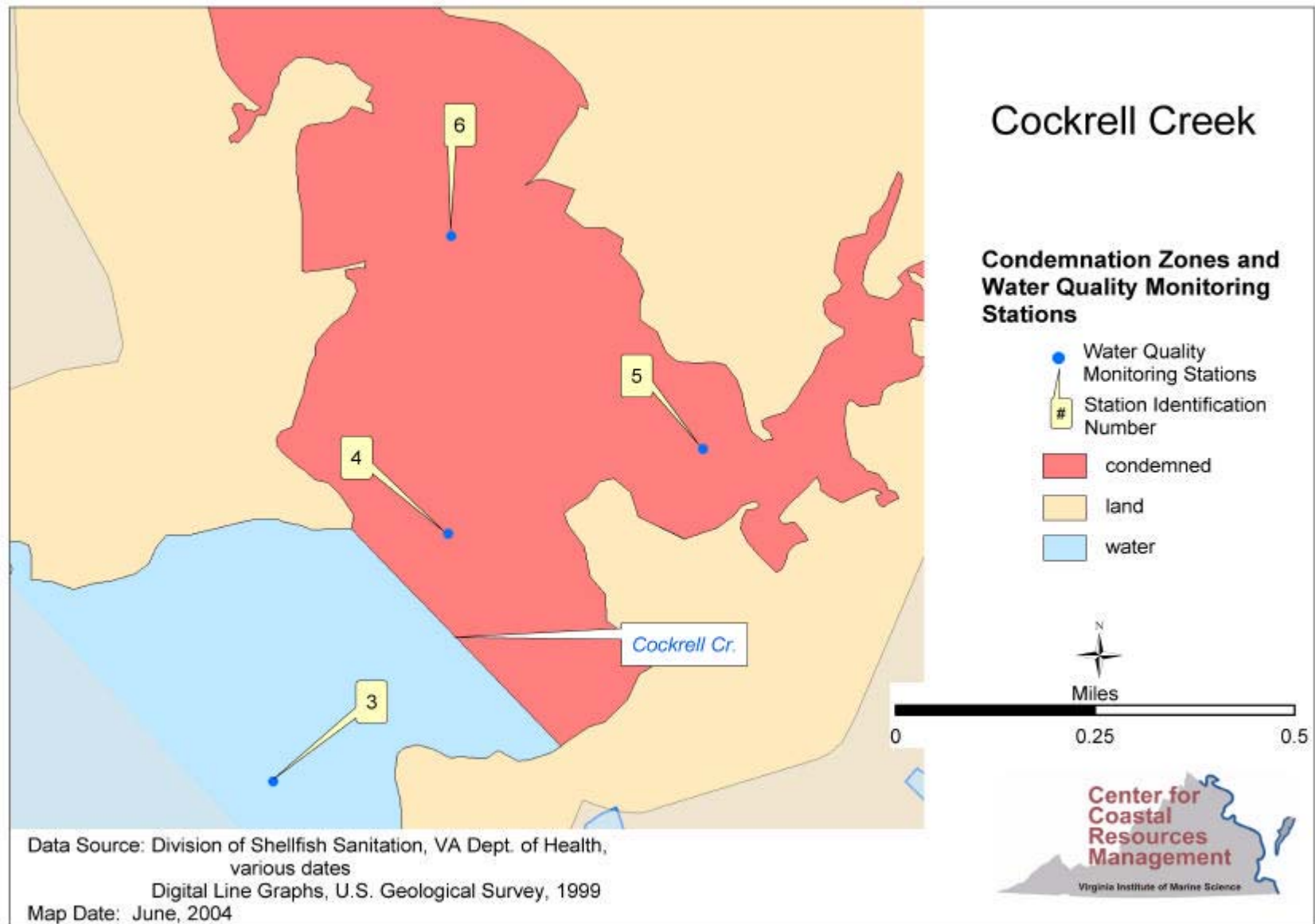
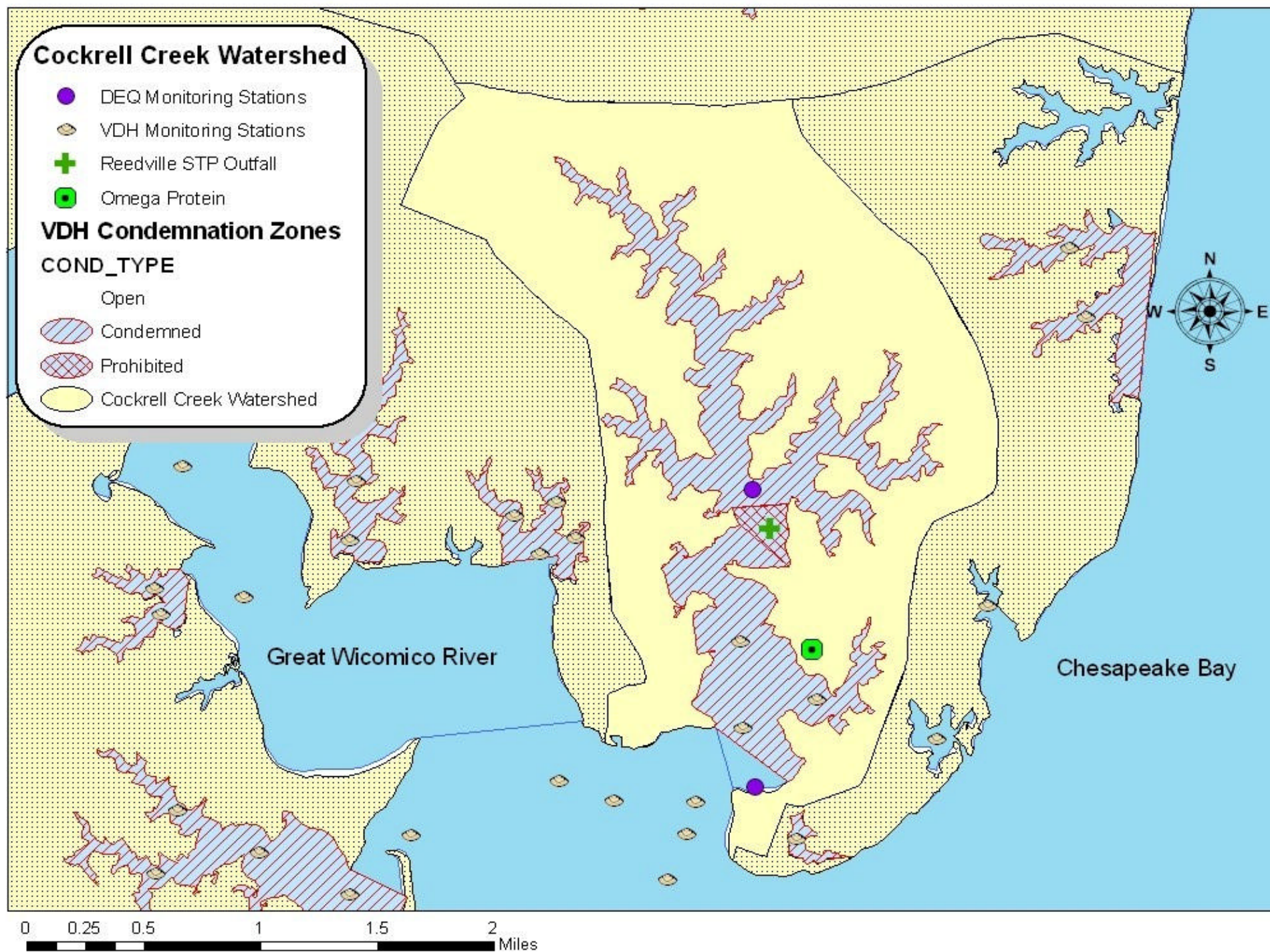


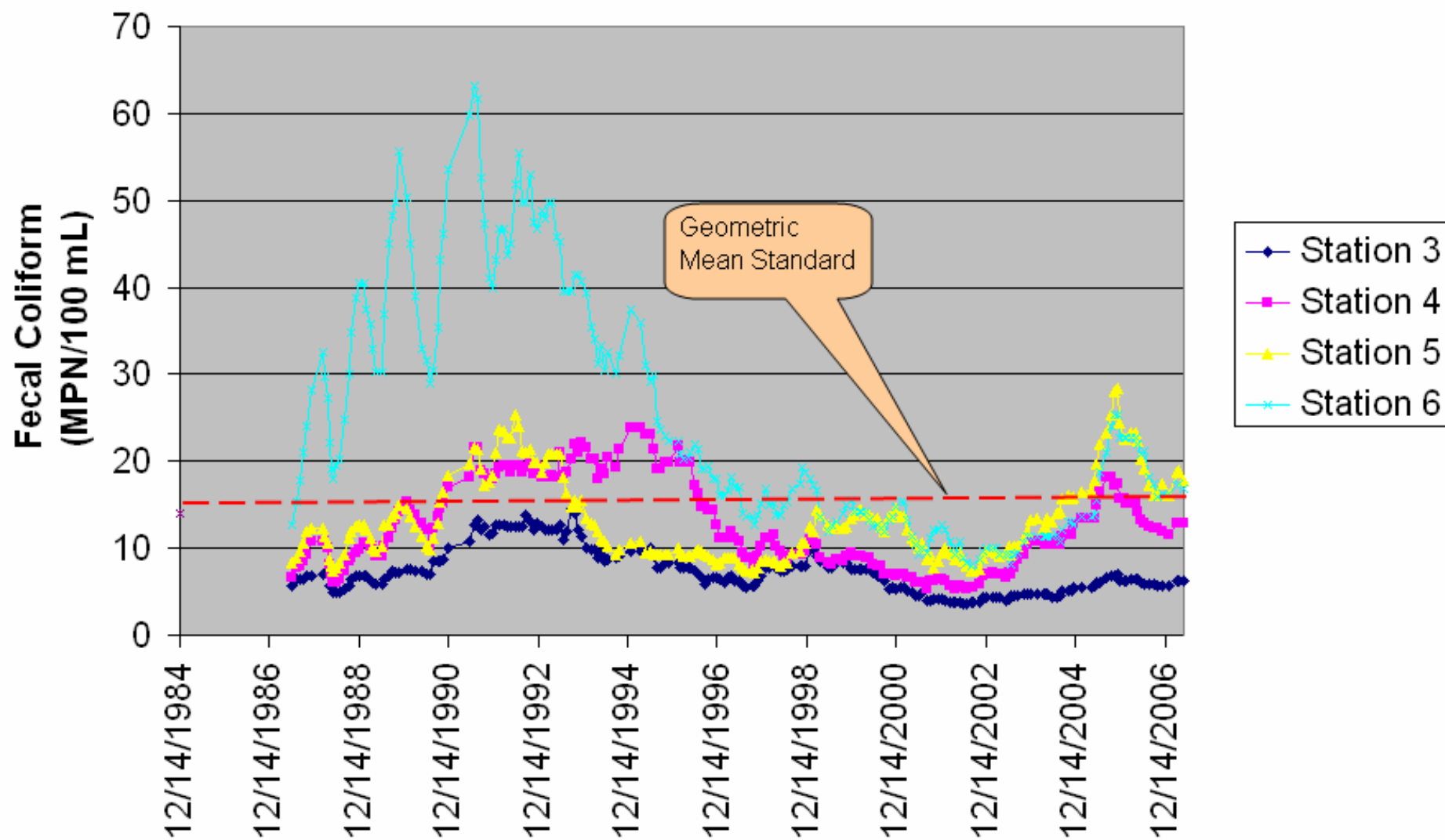
Figure 4.1



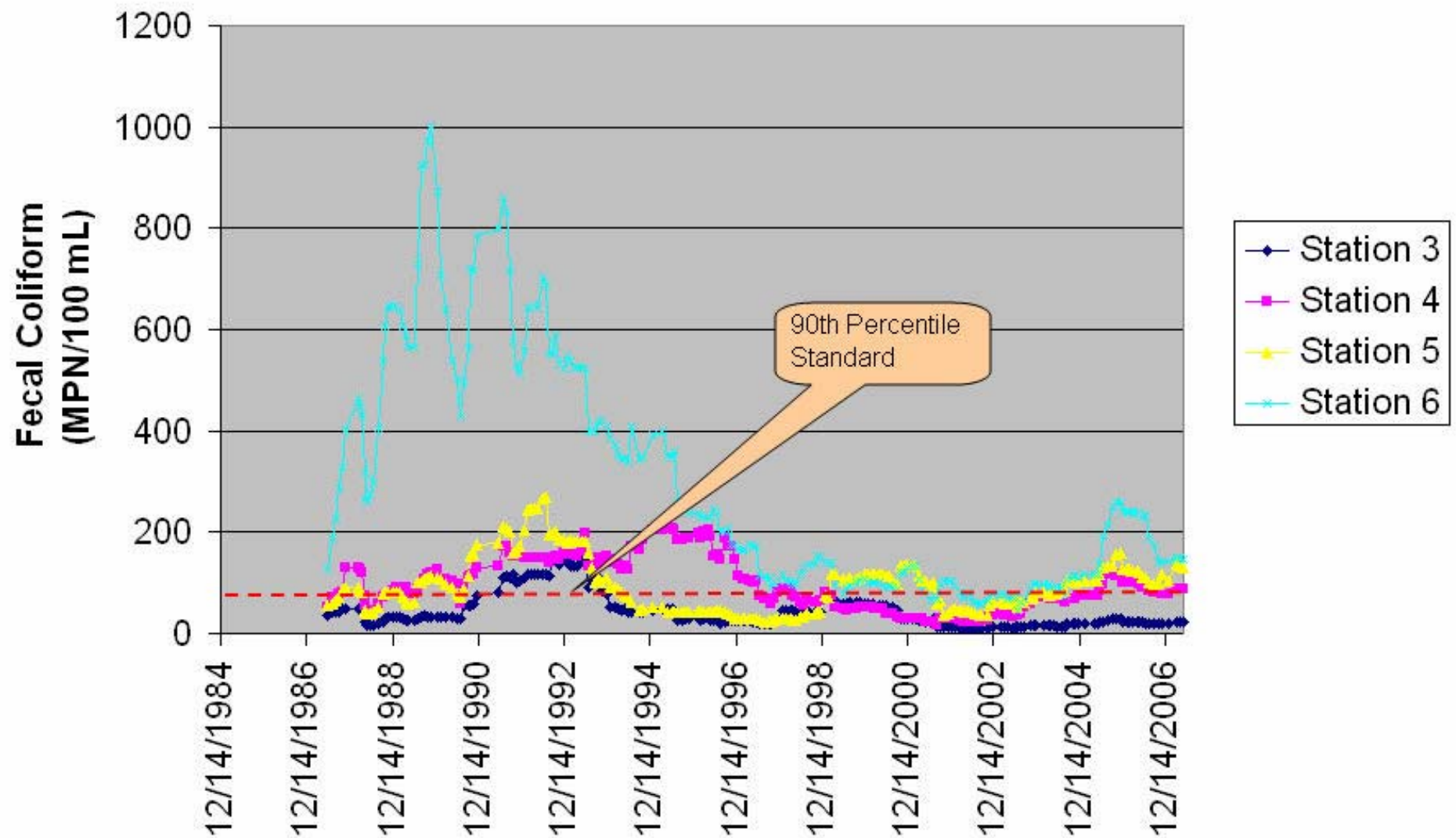
DEQ 2006-2007 Special Study Monitoring Stations



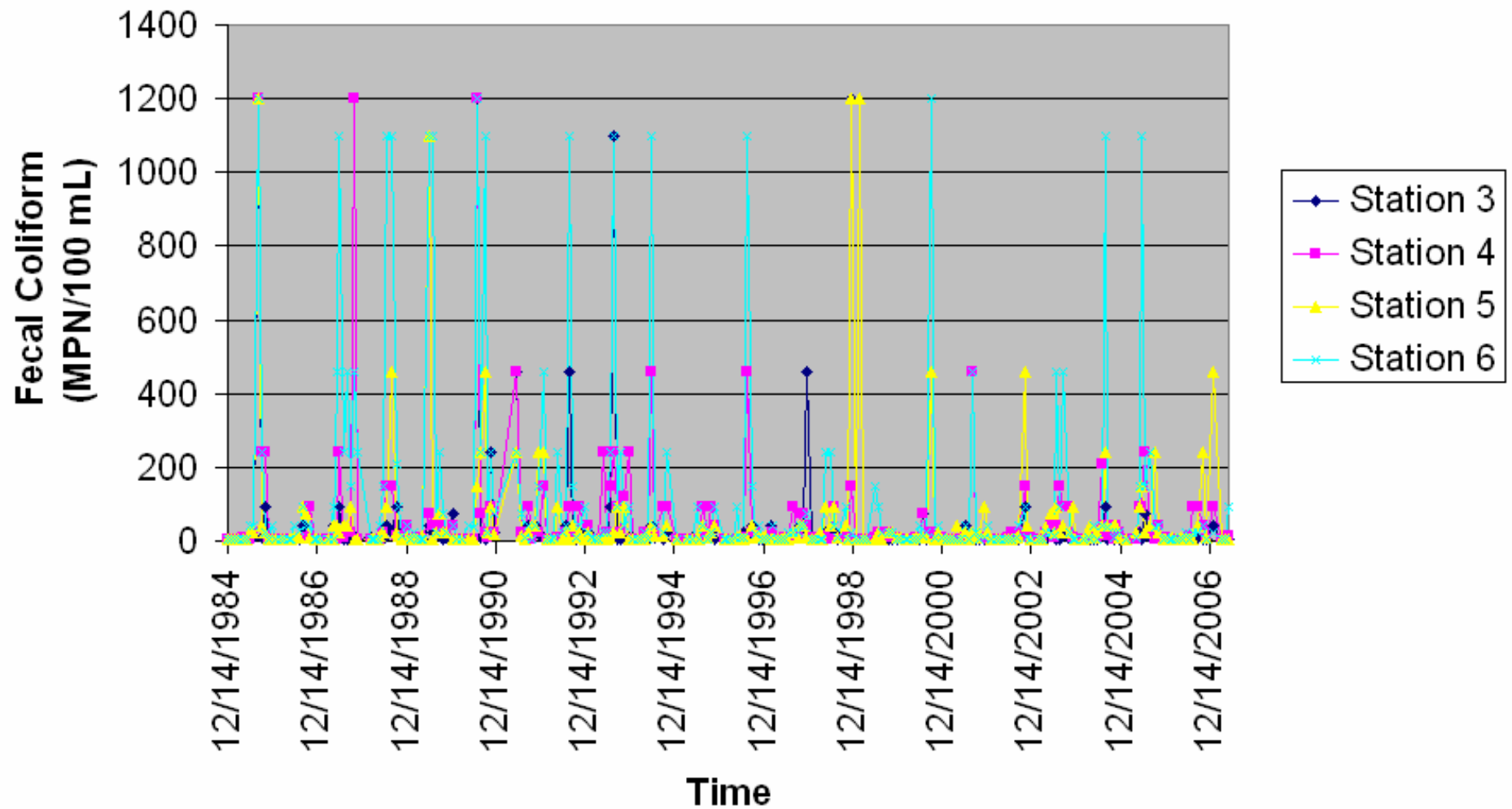
Cockrell Creek - Running Geometric Mean



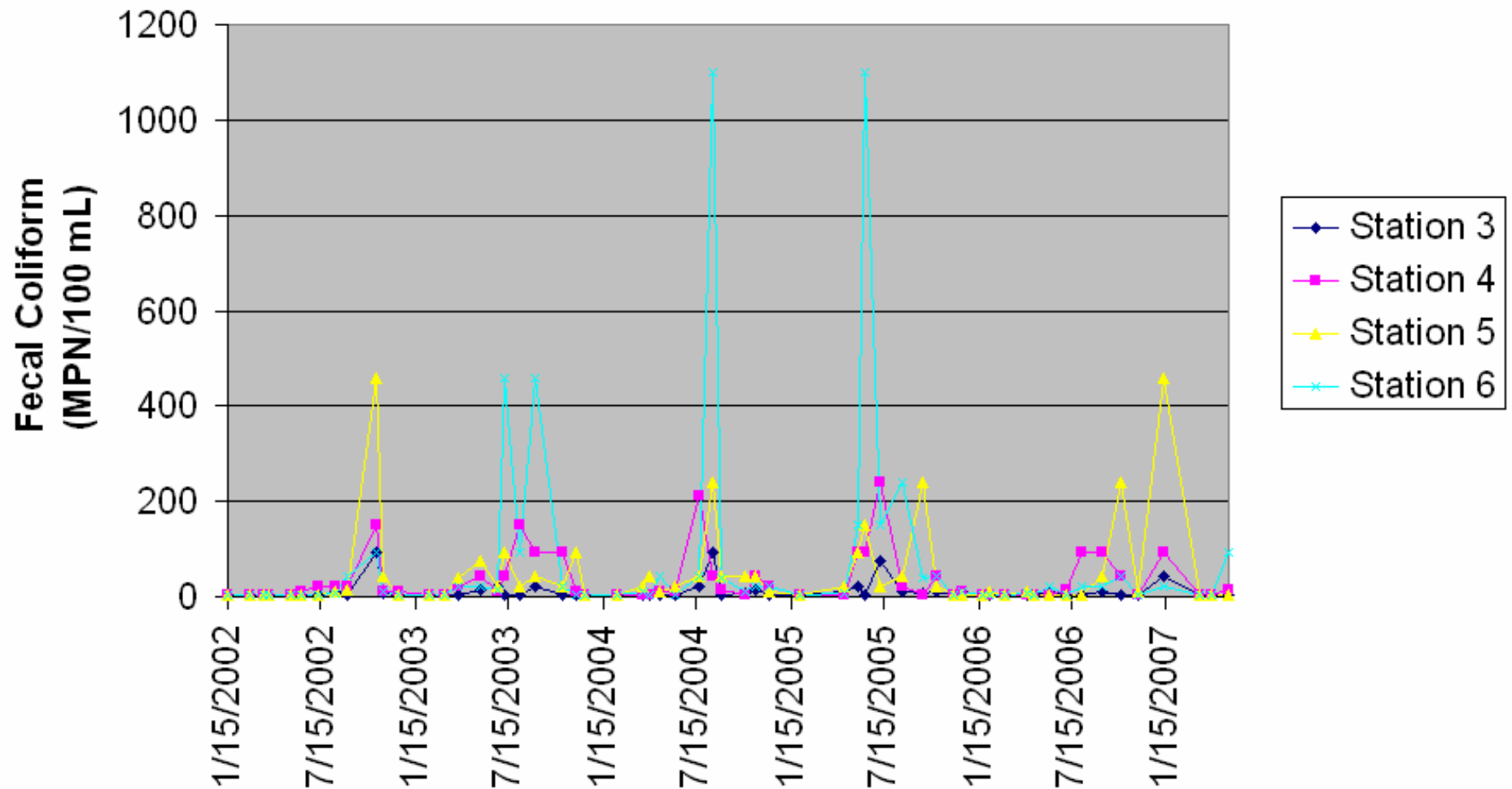
Cockrell Creek - Running 90th Percentile

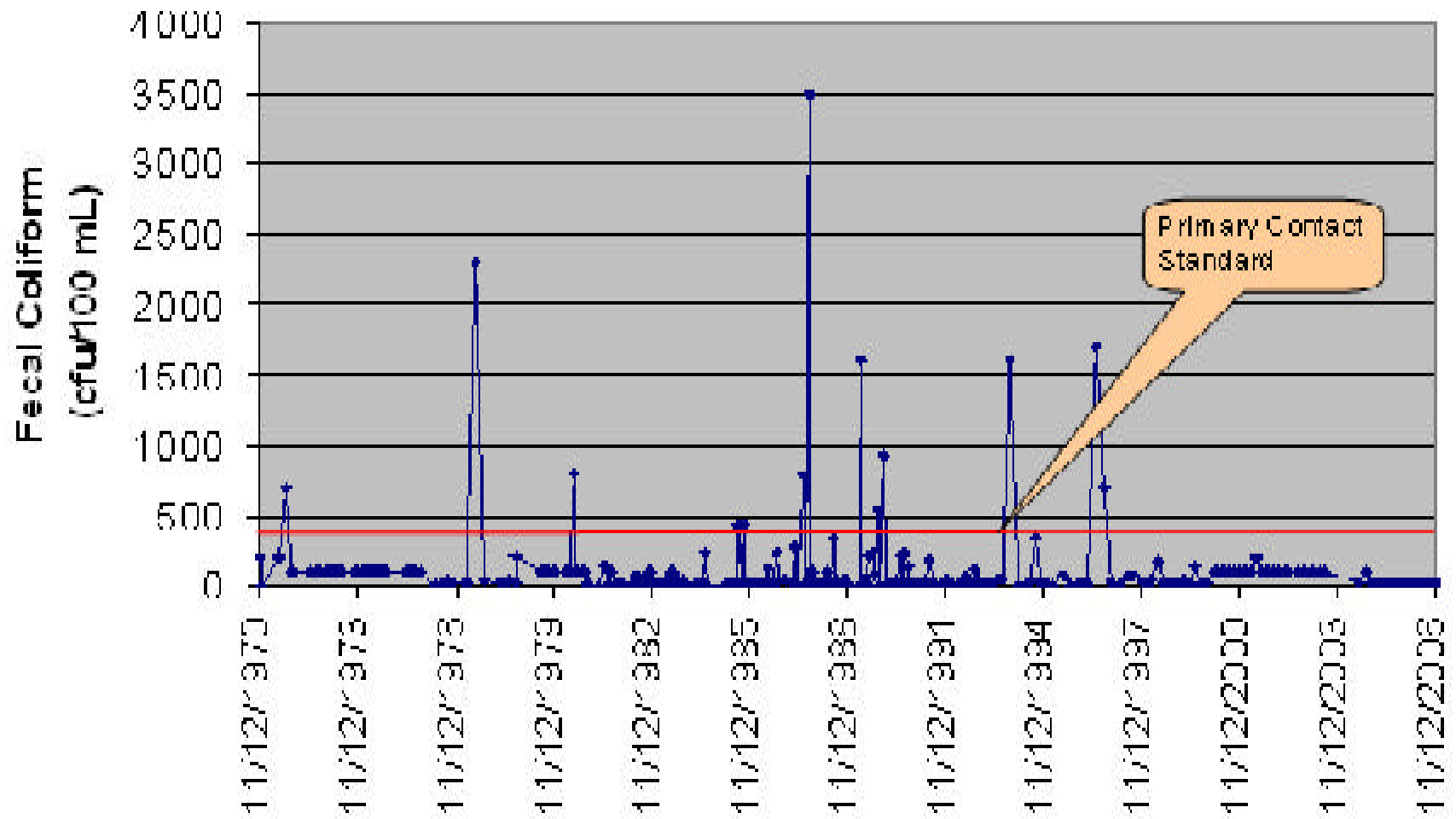


VDH Bacterial Data (Cockrell Creek) 1984-Present



VDH Bacteria Data (Cockrell Creek) 2002 - Present



DEQ Station 7-COC001.61

DEQ Special Study - Fecal coliform Results

Monitoring Station	Station Description	3-Aug-06	20-Sept-06	27-Sept-06	22-Feb-07
7-COC000.86	VDH-DSS Site 12-6	25	<100 U	1000	<10 U
7-COC000.88	SE Corner Omega Pier with Ships	>2000 L	1600	<100 U	<10 U
7-COC000.89	South Side Omega Pier with Ships	520	1000	1000	<10 U
7-COC000.92	West Side Omega Pier with Ships	>2000 L	400	1000	<10 U
7-COC000.95	North Side Omega Pier with Ships	N/A	1700	200	<10 U
7-XAN000.17	VDH-DSS Site 12-5	<25 U	<100 U	4000	<10 U
VA0003867-002	Omega Outfall 002	280	>8000 L	5000	<10 U
VA0003867-995	Omega Outfall 995	>2000 L	10,000	1400	N/A
VA0003867-INT	Omega Water Intake	780	1800	500	<10 U

Fecal Coliform Units = # cfu/100 mL

Bold & Red Bacteria Values = WQ Violations using instantaneous primary contact standard 400 cfu/100mL (fecal coliform)

Bold Bacteria Values = WQ Violations using 90th Percentile shellfish standard - 49 MPN/100mL (fecal coliform)

Com Code L = Off Scale high, actual value greater than the value shown (>max detection limit)

Com Code U = Material analyzed for but not detected (<minimum detection limit)

NA = Not Available

DEQ Special Study – Enterococci Results

Monitoring Station	Station Description	3-Aug-06	20-Sept-06	27-Sept-06	22-Feb-07
7-COC000.86	VDH-DSS Site 12-6	75	110	210	<25 U
7-COC000.88	SE Corner Omega Pier with Ships	1800	3400	680	<25 U
7-COC000.89	South Side Omega Pier with Ships	680	600	240	<25 U
7-COC000.92	West Side Omega Pier with Ships	>2000 L	300	520	<25 U
7-COC000.95	North Side Omega Pier with Ships	N/A	800	1000	<25 U
7-XAN000.17	VDH-DSS Site 12-5	<25 U	<10 U	<10 U	<25 U
VA0003867-002	Omega Outfall 002	300	160	700	<25 U
VA0003867-995	Omega Outfall 995	>2000 L	4600	1400	N/A
VA0003867-INT	Omega Water Intake	>2000 L	>8000 L	>8000 L	<25 U
7-COC001.61	DEQ Ambient Station	N/A	N/A	N/A	25
7-COC000.27	DEQ Ambient Station	N/A	N/A	N/A	75

Enterococci Units = # cfu/100 mL

Bold & Red Bacteria Values = WQ Violations using instantaneous primary contact standard 104 cfu/100mL (Enterococci)

Com Code L = Off Scale high, actual value greater than the value shown (>max detection limit)

Com Code U = Material analyzed for but not detected (<minimum detection limit)

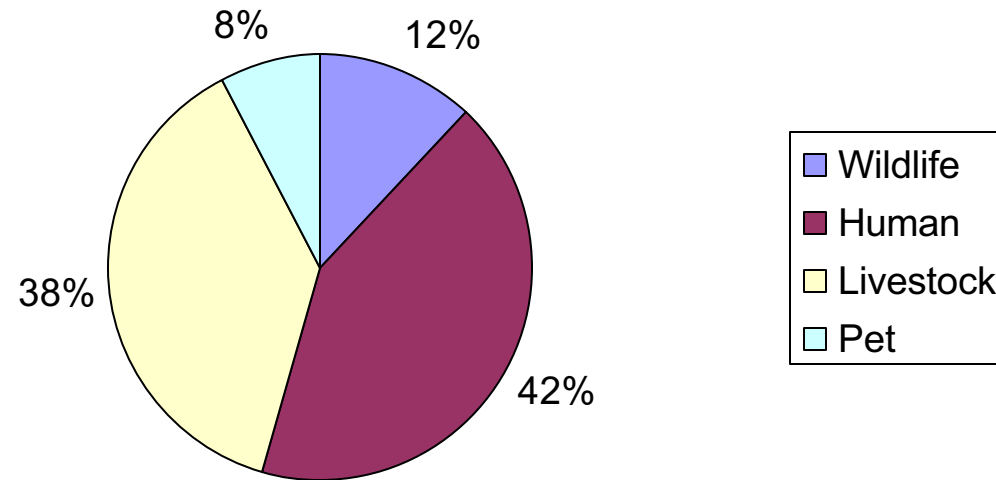
NA = Not Available

Use of BST in the TMDL

- Uses VDH-DSS routine monitoring data to calculate critical fecal count
- Uses 9 supplementary BST samples at selected stations to identify bacteria sources:
- Uses Antibiotic Resistance Analysis as BST method for source load allocation into 4 categories:
 1. **Human**
 2. **Pets**
 3. **Livestock**
 4. **Wildlife**



Weighted Mean Fecal Coliform Contribution By BST Condemned Zone 2A



DEQ Special Study – BST Utilized.

- Significant Human Signature Detected in Omega Protein Intake & Discharges

Water Quality Data Summary

- DEQ station 7-COC001.61 shows long term improvement, attributable to upgrades at the Reedville STP in the 1990's
- VDH-DSS data shows seasonal exceedances of the Shellfish Water Quality Standard
- BST Data shows a significant human source from Omega Protein
- DMR data collected by Omega continues to show high bacterial counts from the treatment lagoon
- BST data from Omega lagoon includes human signature

Water Quality Data Summary

- Many bacterial samples from the Special Study not only exceed the Shellfish Standard, but also the Primary Contact (Swimming) Standard
- Omega DMR data also exceeds the Primary Contact (Swimming) Standard
 - Cockrell Creek will be listed as impaired for Primary Contact in 2008 VA Water Quality Assessment

Additional Information to Support Study

- 2002 Fluorometric Dye Trace Study conducted for Omega Protein
 - Showed Omega vessels influence facility intake
- 2004 VDH-DSS Shoreline Survey
 - All Marinas have Pump-out facilities for boats
 - DEQ station 7-COC001.61 does not show high bacterial values
- 2006-2007 DEQ Special Study
 - Significant human bacterial sources at Omega Intake & discharges
 - No violations of water quality standards during Omega off season

Conclusions

- Omega Protein, Inc. operational activities contribute significantly to the bacterial impairment in Cockrell Creek
- The Omega Fleet is the most probable source of bacteria
 - Ships have Type II Marine Sanitation Devices (MSDs) which discharge at 200 cfu/100mL (>14 times the shellfish standard)
- Omega Lagoon is a source of bacteria
 - DMR data, DEQ sampling, BST Human signature

Conclusions

- Additional non-point sources of bacteria are found within the watershed
- All 4 categories detected with in-stream BST samples

TMDL Calculation

Shellfish Designated Use Load Reductions

Table 5.1. Geometric Mean Analysis of Current Load and Estimated Load Reduction

Condemnation Area	Volume (m ³)	Fecal Coliform (MPN/100ml)	WQ Standard MPN/100 ml	Current Load (MPN/day)	Allowable Load (MPN/day)	Required Reduction (%)
002A Cockrell Creek (VAP-C01E-08)	5102820	37.4	14	1.91E+12	7.14E+11	63%

Table 5.2. 90th Percentile Analysis of Current Load and Estimated Load Reduction

Condemnation Area	Volume (m ³)	Fecal Coliform (MPN/100ml)	WQ Standard MPN/100 ml	Current Load (MPN/day)	Allowable Load (MPN/day)	Required Reduction (%)
002A Cockrell Creek (VAP-C01E-08)	5102820	399.5	49	2.04E+13	2.50E+12	88%

Shellfish TMDL

Reduction and Load Allocation Based Upon 90th Percentile Standard: Cockrell Creek - Growing Area 12

Condemnation Area	Source	BST Allocation % of Total Load	Current Load MPN/ day	Load Allocation MPN/ day	Reduction Needed
002A Cockrell Creek (VAP-C01E-08)	Wildlife	12%	1.60E+12	1.60E+12	0%
	Human	42%	5.59E+12	0.00E+00	100%
	Livestock	38%	5.06E+12	0.00E+00	100%
	Pets	8%	1.06E+12	8.90E+11	16%
	Total	100%	2.04E+13	2.50E+12	88%

Reduction and Waste Load Allocation Based Upon 90th Percentile Standard: Cockrell Creek - Growing Area 12

Condemnation Area	Discharger	Current Load MPN/ day	Load Allocation MPN/ day	Reduction Needed
002A Cockrell Creek (VAP-C01E-08)	Omega Protein (VA0003867)	7.09E+12	9.97E+09	100%

Primary Contact Designated Use TMDL

Table 5.3 Calculations for Recreation Use Impairments in Cockrell Creek

Impaired Area	Volume (m³)	Bacteria Pollutant	Current Load (cfu/day)	Allowable Load (cfu/day)	Required Reduction (%)
002A Cockrell Creek (VAP-C01E-08)	5102820	<i>Enterococci</i>	4.08E+14	5.31E+12	99%

Highest observed ambient exceedence = >8000 cfu at Omega Protein, Inc.

TMDL Summary

Table 5.8. TMDL Summary for the Closure in the Cockrell Creek Watershed (90th percentile)

Condemnation Area	Pollutant Identified	TMDL MPN/day	Waste Load Allocation MPN/day	Load Allocation MPN/day	Margin of Safety
002A Cockrell Creek (VAP-C01E-08)	Fecal Coliform	2.50E+12	9.97 E+09	2.50E+12	Implicit

TMDL Summary for the Recreation Use Impairment in Cockrell Creek

Impaired Water body Segment	Volume (m ³)	Bacteria Pollutant	Load Allocation (cfu/day)	Wasteload Allocation (cfu/day)	TMDL	Margin of Safety
002A Cockrell Creek	5102820	<i>Enterococci</i>	5.29E+12	2.49E+10	5.31E+12	Implicit

Some Suggested Implementation Measures

- Establish No discharge Zone for vessels in Cockrell Creek or Great Wicomico River and tributaries
- More vigorous policing of pump out requirements
- Require higher level of septic treatment in home systems that need replacement and for new homes
- Establish a state cost share/grant program to aid home owners in replacing failing or inadequate systems
- Public education effort regarding animal and pet waste
- Public education effort regarding proper septic tank maintenance

Water Quality Fact

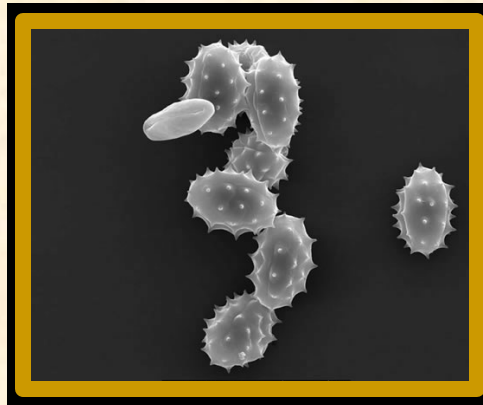
- ❖ “A single weekend boater flushing untreated sewage into our waters produces the same amount of bacterial pollution as 10,000 people whose sewage passes through a treatment plant.”



- ❖ California Department of Boating and Waterways

Water Quality Fact

- ❖ “ A single overboard discharge of human waste can be detected in up to a one square mile area of shallow enclosed water. Contaminants known from human waste include hepatitis, streptococci, fecal coliform and other bacteria.”



- ❖ Florida Department of Environmental Protection

Current No Discharge Zones (NDZs) in VA

- Smith Mountain Lake
- VA Beach Area (Lynnhavan, Broad, & Linkhorn Bays)
- VA Beach Area – Current efforts to establish additional NDZs for entire jurisdiction
- NDZs – possibly the most cost effective BMP to restore local water quality

Next steps

- 30 Day Public Comment Begins
- TMDL will be presented at SWCB meeting
- TMDL will be sent to EPA for approval

Send written comments to:

Mark Alling
4949-A Cox Rd
Glen Allen VA 23060
804/527-5021
msalling@deq.virginia.gov

Omega Protein WLA by Outfall

Table 5.4. Omega Protein, Inc. Fecal Coliform Wasteload Allocation

Omega Protein Outfall	Daily Maximum Design Flow (MGD)	Highest recorded fecal coliform (cfu/100 mL)	Fecal Coliform Existing load	Fecal Coliform Wasteload Allocation
001	4.14	10,000*	1.57E+12	2.19E+09
002	0.481	>8000	1.46E+11	2.55E+08
995	14.2	10,000	5.37E+12	7.52E+09
Total			7.09E+12	9.97E+09

* Bacterial samples not collected at 001 during study. Since source water for 995 & 001 are same, values for 995 were assigned to 001.

Table 5.5. Omega Protein, Inc. Enterococci Wasteload Allocation

Omega Protein Outfall	Daily Maximum Design Flow (MGD)	Highest recorded Enterococci (cfu/100 mL)	Enterococci Existing Loads	Enterococci Wasteload Allocation
001	4.14	4600*	7.21E+11	5.48E+09
002	0.481	700	1.27E+10	6.37E+08
995	14.2	4600	2.47E+12	1.88E+10
Total			3.21E+12	2.49E+10

* Bacterial samples not collected at 001 during study. Since source water for 995 & 001 are same, values for 995 were assigned to 001.